




Parsimonious modelling of spectroscopy data via a Bayesian latent variables approach

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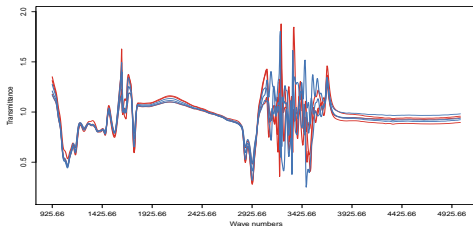
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➤ Cows, Diet & Spectroscopy

- Increasing consumers awareness is changing the food industry
- Cattle **feeding regimen** (pasture vs total mixed ration)
 - pasture diet regarded as more respectful and products as more natural and healthier
 - characterization of the differences implied by different diets on the milk features still overlooked
- **Vibrational Spectroscopy techniques**
 - cheap, rapid and non-disruptive way to collect vast amount of data + widely used in food science

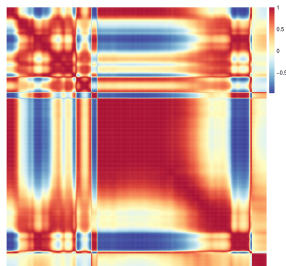
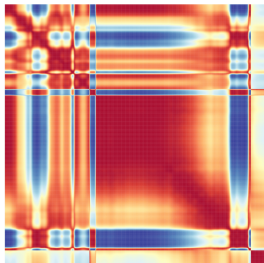


> All that glitters is not gold

- Some challenges:
 - High-dimensionality
 - Peculiar correlation structures
- Aim
 - Parsimonious representation
 - Proper wavelengths relationships reconstruction
 - Characterization of the phenomenon

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Long story short: extract useful knowledge and insights



➤ Latent variable models - Factor Analysis

- We focus on **Factor Analysis** (FA)
 - dimensionality reduction...
 - with a focus on the covariance structure
- Let $X = \{x_1, \dots, x_n\}$ the observed data, FA models $x_i \in \mathbb{R}^p$ as

$$x_i = \Lambda u_i + \varepsilon_i, \quad i = 1, \dots, n$$

$\Lambda \in \mathbb{R}^{p \times K}$ loadings, $u_i \in \mathbb{R}^K$ the scores, K the number of factors

$$x_i \sim \mathcal{N}_p(0, \Sigma = \Lambda \Lambda^T + \Psi)$$

- **Spotlight** on sparse estimation of Λ
→ enhance interpretability, detect uninformaticness

What about **redundancy**?

> Factor Analysis with redundant variables

- **Idea** search for variable clustering structures
→ modify standard FA by allowing some variables to be mapped by means of the same loadings
- Proposed model

$$x_i = Z\Lambda_c u_i + \varepsilon_i \quad i = 1, \dots, n$$

- $Z \in \mathbb{R}^{p \times G}$ latent *allocation matrix*, $z_{jg} = 1$ if j -th variable belongs to g -th cluster, G the number of variable clusters
- $\Lambda_c \in \mathbb{R}^{G \times K}$ cluster specific loadings
- Number of loadings: $(p \times K) \rightarrow (G \times K)$

➤ Additional thoughts

- In matrix form

$$\begin{pmatrix} x_{i,1} \\ \vdots \\ x_{i,p} \end{pmatrix} = \begin{pmatrix} z_{1,1} & \cdots & z_{1,G} \\ \vdots & \ddots & \vdots \\ z_{p,1} & \cdots & z_{p,G} \end{pmatrix} \begin{pmatrix} \lambda_{1,1}^c & \cdots & \lambda_{1,K}^c \\ \vdots & \ddots & \vdots \\ \lambda_{G,1}^c & \cdots & \lambda_{G,K}^c \end{pmatrix} \begin{pmatrix} u_{i,1} \\ \vdots \\ u_{i,K} \end{pmatrix} + \begin{pmatrix} \varepsilon_{i,1} \\ \vdots \\ \varepsilon_{i,p} \end{pmatrix}$$

- Uniformitiveness can be detected by forcing a row of Λ_c to be equal to a zero vector
- Denoting with $\tilde{\Lambda} = Z\Lambda_c$ and adapting distributional assumptions we obtain

$$(x_i|Z) \sim \mathcal{N}_p(0, \tilde{\Sigma} = \tilde{\Lambda}\tilde{\Lambda}^T + \Psi)$$

> Model estimation - Priors & Algorithm

- Bayesian estimation procedure is adopted
 - Standard prior distribution for Λ_c, u_i, Ψ
 - Prior on the allocation matrix \rightarrow **Product Partition Model**
Correspondence between Z and $\mathbf{c} = \{C_1, \dots, C_G\}$

$$\pi(\mathbf{c}) \propto \prod_{g=1}^G \rho(C_g) = \alpha_z^G \prod_{g=1}^G (|C_g| - 1)!$$

- Conjugate nature of the priors allows a Gibbs updating scheme
- MH-step to sample Z via modification of the **allocation sampler**
 - Idea** a single move attempts to reallocate a block of variables from one group to another
 \rightarrow Bigger moves, faster exploration
 - Modification aim to propose moves involving *close* clusters

> Model selection

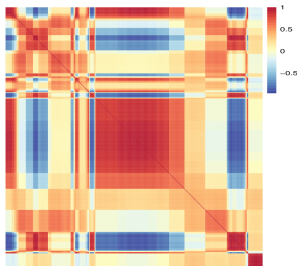
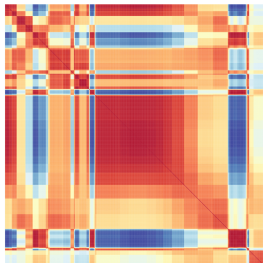
- Number of factors K and number of variable clusters G
- Possible solutions:
 - Information criteria (AICM, BICM, BIC-MCMC...)
 - Nonparametric fashion \rightarrow infinite groups/factors
- **Proposal** \rightarrow ad-hoc initialization strategy
 - Idea: multi-step procedure to mimick $\tilde{\Lambda}$ structure via standard FA and model-based clustering strategy
 - Output: $(K_{\text{init}}, G_{\text{init}})$ initial guess to be used as the starting for a local search
 - Rationale: avoid intensive global search since the focus is on covariance reconstruction

➤ Dairy diet MIRS data

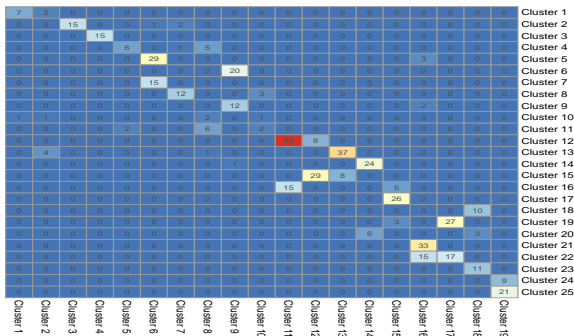
- Milk samples from 120 cows collected weekly during summer months (from 2015 to 2017)
- Two different feeding regimens considered:
 - Pasture, cows maintained outdoors on perennial ryegrass and white clover
 - Total mixed ration (TMR), cows maintained indoors, nutrients combined in a single mix
- $n = 4320$ milk samples, $n_P = 2391$ $n_T = 1389$
 $p = 1060$ wavelengths in the mid-infrared region
→ water-absorption spectral regions removed thus $p^* = 533$

> Some results - MIRS data

- Initialization strategy suggests
($K_P = 4, G_P = 25$) , ($K_T = 3, G_T = 19$)
 - Good reconstruction of the sample correlations
 - Blocky structure as a byproduct of the clustering mechanism
- ↓
- useful to highlight differences between spectral regions



➤ Some results - MIRS data



- Quite strong agreement between the two partitions ($ARI = 0.65$)
→ signal about real and non-spurious clustering structures
- Pronounced redundancy, grouping may be used to build new variables or for cluster-specific predictive analyses
- **Interpretability** → subject-matter knowledge is needed

➤ Concluding remarks

- The proposed method provides parsimonious summaries of high dimensional data with highly correlated variables
→ not only spectroscopic data
- Richer insights with respect to standard FA thanks to the variable clustering mechanism
- **Directions and open questions**
 - It might serve as a building block for classification tools, easy to embed in a MFA framework
 - Different choices for the priors?
 - shrinkage priors
 - exploit some *spatial* information when specifying the cohesion function

> Some references

Check the paper out on arXiv

<https://arxiv.org/pdf/2101.12499.pdf>

Other references

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